

1 2 3 4 5 6 7 8 9 10 11 12

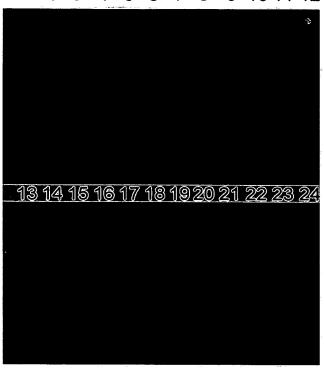


FIG.2A

7 8 9 10 11 12

13 14 15 16 17 18 19 20 21 22 23 24

FIG.2B

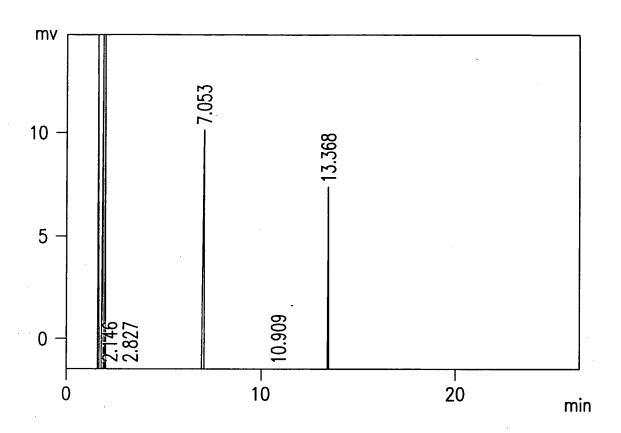
	== <u>_</u>	1H 1 7-51
SP 6	Hind III Sph - Hinc III Acc - Sal - Sal - Sau 3A	
	· ·	0.5 kb
	Sal	1.0 kb
		0.4 kb
오	Pvu	
pSa240	, 	1.0 kb
	Stu	
		1.0 kb
	Sal	1.0 kb
7.1	EcoR   Sac   Kpn   Ava   Sma   BamHI / Sau3A	

FIG.7

## Sa240 DNA sequence (4,826 bp)

gatcggcggc	cggtcggcgg	tgctggccgc	ggtgaccctg	ggggcgctgg	ccgctccggc	60
			cgcggaggcg			120
			cgcggtggcc			180
			cgcggcgctg			240
gctgggcaag	ctgcgcctgc	cgttgccggc	cgccgtggtg	ctggcccagt	ggccgctgct	300
gttctgggcc	tgggccgtgg	gcgcaccggc	gccggtggtc	gggtgggcgc	tgctggccac	360
cgcggtgctg	gacggggcga	tcgccctgtg	gggcaagggc	gccggggtgc	gggtcacggc	420
gtgcgtcggt	ggagcggtga	tgggcttctc	ggccctgatg	gtgggcctgg	cgctgtccct	480
gacggccccg	gggccgctcg	gggcggtggc	tccgggcgtg	ctgctgctga	cggcctcggc	540
ggcggccgtg	gccggggcgt	ggcgcgcgcc	gaagggtttc	gcgcggacgg	gtggtgcggt	600
ggcggggctc	gcggcggtgg	cggccgtcgg	cggcgtaccg	gcggcggcgc	tcccggcggg	660
ctggcgggtg	ctcgcgtacc	tgctgtgcgg	tctcgcgttg	acggcggtcg	tccgttcccg	720
gctgccgggc	cacgccgcgc	gcggggtact	ggcggcgtcg	ggggcggtgg	tggccggcgc	780
gctggtgtgg	gcgctgccgc	cgctcgcggc	ggtgctgctg	gggccggtga	cggtgctgtc	840
ggacgtgtgg	gcggggacgc	cggacggctt	ccggtccgcg	ctggggtcga	cgctgccctg	900
			cgcgctggtg			960
gcgtaaccgg	aggtggccgt	cggtcgtccg	gctccggcgc	cgttggccgg	tccttctggc	1020
			cgggagcccc			1080
cggccggggg	cgctgcgccg	tggcccggct	ggtccggctg	gtccggccgg	cccggtgcgg	1140
gggccggtgg	tcgcgggcgg	ccttccgcgg	cgacgctgcg	cggggtcgtc	ggcgcgggcg	1200
			tggccggcgc			1260
			tgggcgtcct			1320
gtggcggcgc	cgagcggggc	gcgacggcga	tgccggtgac	cgctctggtg	gcttcggtgg	1380
ccggggcggt	gagcgccggg	ctgctgtcgc	tggcgtccga	gggggcctcg	tacgcggtgt	1440
tcggcgcgct	ggcggcgctg	ttcgccgggg	ccgctctgcg	ggcgggcgcc	ggggtgccgc	1500
			ggggcaccgt			1560
			ccccgctgat			1620
cggtgctgct	cggggcacga	ctgcggcgga	acccggtggc	cttgcccgtg	gagctgacgg	1680
gagcgctggg	cgcgctcgtc	gccgtggggc	tcgcggtgtc	cgacgcgccg	ttcctggccc	1740
			cggggacggc			1800
cggtggcggg	ctacctggcg	gcgacgctgt	tcgtgctggc	cacgtgggtg	cggctggcgg	1860
cctcggaggt	gtcgttcccg	gaggcgtaca	cgctgccggt	gacggtgccc	gcgctgctgg	1920
			aggcctcgtc			1980
			cggtcgcctg			2040
			tgatcaccct			2100
			tgctggcact			2160
			tccccgctg			2220
			acgagcagcg			2280
			ccgtgcccgg			2340
			ggcaacgcag			2400

ggtgggcgat	actgggttcg	aaccagtgac	ctcttcggtg	tgaacgaagc	gctctcccac	2460
	gcccgggcgc					2520
	actcgctgat				•	2580
ccggccagca	ccgccatgat	cacgagcccg	agcgtggtga	ggatgatgtt	gcgccgccgg	2640
accttgggat	cgagggcccg	ctgcgccgct	tcggtgacct	tgcgcttggt	ccagcgcagc	2700
	cccagacgaa					2760
	ccggcagcac					2820
	cgacctgcca					2880
	ccagcgcgcg					2940
gcctgctcgg	cgaccttgct	ccgctcgtca	ctctccgcgt	tcatgaagct	caacttaccc	3000
gacctgtctc	cgtcactgga	atgggcgcat	aactcaaagt	tacacgccgc	tgagcggggg	3060
acccgaagcg	tcacaaatgg	gtcagagggg	tttacaacgc	caccgtaggt	ggcatgtcga	3120
tttcgccgac	gtgcgaatcc	ccgagcgcac	actgagcgaa	aggccctggc	gcttatgaac	3180
accacggtca	gctgcgagct	gcacctgcgc	ctcgttgtgt	cgagcgagtc	ctcactgcct	3240
gtacccgcgg	gcctgcggta	tgacacggcc	gatccctatg	ccgtgcacgc	caccttccac	3300
accggagcgg	aggagacggt	cgaatgggta	ttcgcccgcg	acctccttgc	cgaggggctg	3360
caccggccca	ccggcaccgg	agacgtccgc	gtctggccat	ctcgtagtca	cggtcaaggc	3420
gtcgtatgca	tcgccctgag	ctccccagag	ggagaagccc	tgctcgaagc	cccggcgcgg	3480
gccctggagt	cgttcctgaa	gaggaccgac	gccgcggttc	cgcccggcac	cgagcatcgt	3540
	tcgacacgga					3600
gccgctctac	gccgtccgac	tcggggcgac	ggcgtcgtgc	tgacaaccgc	atagggcaga	3660
	gtcgtcgcgg					3720
	ggtccgttcc					3780
cgagccagta	gagtcagccg	ccatcggcag	gcgcccgccc	gccggaaggc	cagggagcga	3840
agcgtgctga	tccctcacga	cacccggatc	gccctcgacg	cggtggtcga	tctggtgaac	3900
	agagcgagcc					3960
gaggacggtc	tccccgacat	cgccgcgctg	tacgccttcg	cggagcgcca	tctcatcagc	4020
ggggtcggca	ccctcggcga	gaaggacctc	ggcgccgtgc	gcgacgtccg	ggcccgcttc	4080
	tcgcggcgcc					4140
	ggaccacccc					4200
	cggacgcctc					4260
	tcgtggcggg					4320
	tcgtcgacct					4380
	ggctccacgt					4440
	ggtggcgcga					4500
	ccgatcaccg					4560
	gcctcttcgg					4620
	tcatgacgca					4680
	gtcccgtggc					4740
	gcccggaacg		gggccggccg	acgcctcgga	cgccgcgctt	4800
ctcagatgcc	gtgcttcttg	aggatc				4826



\*\*\* Peak Report \*\*\*

KNO	TIME	AREA	HEIGHT	MK	IDNO	CONC	NAME
1	1.742	1090524	630697	٧		16.3090	
2	1.803	2759591	1055567	VE		41.2702	
3	1.931	841734	440026	٧		12.5883	
4	1.989	1899444	1050304	SVE		28.4066	
5	2.146	1222	1166	T		0.0183	
6	2.827	1691	1181	V		0.0253	
7	7.053	56479	14295			0.8447	•
8	10.909	1014	280			0.0152	
9	13.388	34942	11651			0.5226	
		6686640	3205166		<del></del>	100.0000	

FIG.10

